

1647

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/494,585

DATE: 08/15/2001
 TIME: 12:29:51

Input Set : A:\Cura57U1.APP
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3 <110> APPLICANT: Shimkets, Richard A
 4 Prayaga, Sudhirdas K
 6 <120> TITLE OF INVENTION: Novel Fibroblast Growth Factor and Nucleic Acids
 7 Encoding Same
 9 <130> FILE REFERENCE: 15966-557 FGF-CX
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/494,585
 C--> 12 <141> CURRENT FILING DATE: 2001-01-31
 14 <150> PRIOR APPLICATION NUMBER: USSN 60/145,899
 15 <151> PRIOR FILING DATE: 1999-07-27
 17 <160> NUMBER OF SEQ ID NOS: 13
 19 <170> SOFTWARE: PatentIn Ver. 2.1
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 35 ttg ggc cag cag gtg ggt tcg cat ttc ctg ttg cct cct gcc ggg gag 96
 36 Leu Gly Gln Gln Val Gly Ser His Phe Leu Leu Pro Pro Ala Gly Glu
 37 20 25 30
 39 cgg ccg ccg ctg ctg ggc gag cgc agg agc gcg gcg gag cgg agc gcg 144
 40 Arg Pro Pro Leu Leu Gly Glu Arg Arg Ser Ala Ala Glu Arg Ser Ala
 41 35 40 45
 43 cgc ggc ggg ccg ggg gct gcg cag ctg gcg cac ctg cac ggc atc ctg 192
 44 Arg Gly Gly Pro Gly Ala Ala Gln Leu Ala His Leu His Gly Ile Leu
 45 50 55 60
 47 cgc cgc cgg cag ctc tat tgc cgc acc ggc ttc cac ctg cag atc ctg 240
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 49 65 70 75 80
 51 ccc gac ggc agc gtg cag ggc acc cgg cag gac cac agc ctc ttc ggt 288
 52 Pro Asp Gly Ser Val Gln Gly Thr Arg Gln Asp His Ser Leu Phe Gly
 53 85 90 95
 55 atc ttg gaa ttc atc agt gtg gca gtg gga ctg gtc agt att aga ggt 336
 56 Ile Leu Glu Phe Ile Ser Val Ala Val Gly Leu Val Ser Ile Arg Gly
 57 100 105 110
 59 gtg gac agt ggt ctc tat ctt gga atg aat gac aaa gga gaa ctc tat 384
 60 Val Asp Ser Gly Leu Tyr Leu Gly Met Asn Asp Lys Gly Glu Leu Tyr
 61 115 120 125
 63 gga tca gag aaa ctt act tcc gaa tgc atc ttt agg gag cag ttt gaa 432
 64 Gly Ser Glu Lys Leu Thr Ser Glu Cys Ile Phe Arg Glu Gln Phe Glu
 65 130 135 140
 67 gag aac tgg tat aac acc tat tca tct aac ata tat aaa cat gga gac 480

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68 Glu Asn Trp Tyr Asn Thr Tyr Ser Ser Asn Ile Tyr Lys His Gly Asp
69 145 150 155 160
71 act ggc cgc agg tat ttt gtg gca ctt aac aaa gac gga act cca aga 528
72 Thr Gly Arg Arg Tyr Phe Val Ala Leu Asn Lys Asp Gly Thr Pro Arg
73 165 170 175
75 gat ggc gcc agg tcc aag agg cat cag aaa ttt aca cat ttc tta cct 576
76 Asp Gly Ala Arg Ser Lys Arg His Gln Lys Phe Thr His Phe Leu Pro
77 180 185 190
79 aga cca gtg gat cca gaa aga gtt cca gaa ttg tac aag gac cta ctg 624
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100 Arg Pro Pro Leu Leu Gly Glu Arg Arg Ser Ala Ala Glu Arg Ser Ala
101 35 40 45
103 Arg Gly Gly Pro Gly Ala Ala Gln Leu Ala His Leu His Gly Ile Leu
104 50 55 60
106 Arg Arg Arg Gln Leu Tyr Cys Arg Thr Gly Phe His Leu Gln Ile Leu
107 65 70 75 80
109 Pro Asp Gly Ser Val Gln Gly Thr Arg Gln Asp His Ser Leu Phe Gly
110 85 90 95
112 Ile Leu Glu Phe Ile Ser Val Ala Val Gly Leu Val Ser Ile Arg Gly
113 100 105 110
115 Val Asp Ser Gly Leu Tyr Leu Gly Met Asn Asp Lys Gly Glu Leu Tyr
116 115 120 125
118 Gly Ser Glu Lys Leu Thr Ser Glu Cys Ile Phe Arg Glu Gln Phe Glu
119 130 135 140
121 Glu Asn Trp Tyr Asn Thr Tyr Ser Ser Asn Ile Tyr Lys His Gly Asp
122 145 150 155 160
124 Thr Gly Arg Arg Tyr Phe Val Ala Leu Asn Lys Asp Gly Thr Pro Arg
125 165 170 175
127 Asp Gly Ala Arg Ser Lys Arg His Gln Lys Phe Thr His Phe Leu Pro
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133 Met Tyr Thr
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139 <211> LENGTH: 38

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141 <213> ORGANISM: Artificial Sequence
143 <220> FEATURE:
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170 ttctggaatt tatcagtata gcagtgggcc tggtcagcat tcgaggcgtg gacagtggac 180
171 tctacctcgg gatgaatgag aagggggagc tgtatggatc agaaaaacta acccaagagt 240
172 gtgtattcag agaacagttc gaagaaaact ggtataatac gtactcgtca aacctatata 300
173 agcacgtgga cactggaagg cgatactatg ttgcattaaa taaagatggg accccgagag 360
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187 cagcggcggc cgctccccgg caggaggcaa caggaaatgc gaaccacct gctggcccaa 240
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199 tttgttaagt gccacaaaat acctgcggcc agtgtctcca tgtttatata tgtagatga 180
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201 ctccgtgaaag agaga 255
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227 35 40 45
229 Pro Ala Val Thr Asp Leu Asp His Leu Lys Gly Ile Leu Arg Arg Arg
230 50 55 60
232 Gln Leu Tyr Cys Arg Thr Gly Phe His Leu Glu Ile Phe Pro Asn Gly
233 65 70 75 80
235 Thr Ile Gln Gly Thr Arg Lys Asp His Ser Arg Phe Gly Ile Leu Glu
236 85 90 95
238 Phe Ile Ser Ile Ala Val Gly Leu Val Ser Ile Arg Gly Val Asp Ser
239 100 105 110
241 Gly Leu Tyr Leu Gly Met Asn Glu Lys Gly Glu Leu Tyr Gly Ser Glu
242 115 120 125
244 Lys Leu Thr Gln Glu Cys Val Phe Arg Glu Gln Phe Glu Glu Asn Trp
245 130 135 140
247 Tyr Asn Thr Tyr Ser Ser Asn Leu Tyr Lys His Val Asp Thr Gly Arg
248 145 150 155 160
250 Arg Tyr Tyr Val Ala Leu Asn Lys Asp Gly Thr Pro Arg Glu Gly Thr
251 165 170 175
253 Arg Thr Lys Arg His Gln Lys Phe Thr His Phe Leu Pro Arg Pro Val
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275 Leu Asn Asp His Leu Gly Gln Ser Glu Ala Gly Gly Leu Pro Arg Gly
276 35 40 45
278 Pro Ala Val Thr Asp Leu Asp His Leu Lys Gly Ile Leu Arg Arg Arg
279 50 55 60

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281 Gln Leu Tyr Cys Arg Thr Gly Phe His Leu Glu Ile Phe Pro Asn Gly
282 65 70 75 80
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285 85 90 95
287 Phe Ile Ser Ile Ala Val Gly Leu Val Ser Ile Arg Gly Val Asp Ser
288 100 105 110
290 Gly Leu Tyr Leu Gly Met Asn Glu Lys Gly Glu Leu Tyr Gly Ser Glu
291 115 120 125
293 Lys Leu Thr Gln Glu Cys Val Phe Arg Glu Gln Phe Glu Glu Asn Trp
294 130 135 140
296 Tyr Asn Thr Tyr Ser Ser Asn Leu Tyr Lys His Val Asp Thr Gly Arg
297 145 150 155 160
299 Arg Tyr Tyr Val Ala Leu Asn Lys Asp Gly Thr Pro Arg Glu Gly Thr
300 165 170 175
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325 35 40 45
327 Pro Ala Val Thr Asp Leu Asp His Leu Lys Gly Ile Leu Arg Arg Arg
328 50 55 60
330 Gln Leu Tyr Cys Arg Thr Gly Phe His Leu Glu Ile Phe Pro Asn Gly
331 65 70 75 80
333 Thr Ile Gln Gly Thr Arg Lys Asp His Ser Arg Phe Gly Ile Leu Glu
334 85 90 95
336 Phe Ile Ser Ile Ala Val Gly Leu Val Ser Ile Arg Gly Val Asp Ser
337 100 105 110
339 Gly Leu Tyr Leu Gly Met Asn Glu Lys Gly Glu Leu Tyr Gly Ser Glu
340 115 120 125
342 Lys Leu Thr Gln Glu Cys Val Phe Arg Glu Gln Phe Glu Glu Asn Trp
343 130 135 140
345 Tyr Asn Thr Tyr Ser Ser Asn Leu Tyr Lys His Val Asp Thr Gly Arg
346 145 150 155 160
348 Arg Tyr Tyr Val Ala Leu Asn Lys Asp Gly Thr Pro Arg Glu Gly Thr
349 165 170 175
351 Arg Thr Lys Arg His Gln Lys Phe Thr His Phe Leu Pro Arg Pro Val
352 180 185 190
354 Asp Pro Asp Lys Val Pro Glu Leu Tyr Lys Asp Ile Leu Ser Gln Ser
355 195 200 205

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VERIFICATION SUMMARY

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TIME: 12:29:52

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Output Set: N:\CRF3\08152001\I494585.raw

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L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date